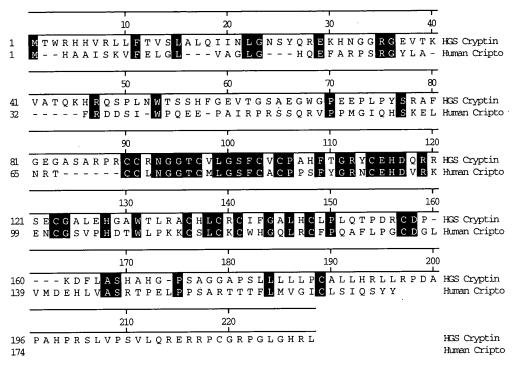
FIG. 1

ATG Met 1	ACC Thr	TGG Trp	AGG Arg	CAC His 5	CAT His	GTC Val	AGG Arg	CTT Leu	CTG Leu 10	TTT Phe	ACG Thr	GTC Val	AGT Ser	TTG Leu 15	GCA Ala		48
TTA Leu	CAG Gln	ATC Ile	ATC Ile 20	AAT Asn	TTG Leu	GGA Gly	AAC Asn	AGC Ser 25	TAT Tyr	CAA Gln	AGA Arg	GAG Glu	AAA Lys 30	CAT His	AAC Asn		96
GGC Gly	GGT Gly	AGA Arg 35	GGG Gly	GAA Glu	GTC Val	ACC Thr	AAG Lys 40	GTT Val	GCC Ala	ACT Thr	CAG Gln	AAG Lys 45	CAC His	CGA Arg	CAG Gln	1	.44
TCA Ser	CCG Pro 50	CTT Leu	AAC Asn	TGG Trp	ACC Thr	TCC Ser 55	AGT Ser	CAT His	TTC Phe	GGA Gly	GAG Glu 60	GTG Val	ACT Thr	GGG Gly	AGC Ser	1	.92
GCC Ala 65	GAG Glu	GGC Gly	TGG Trp	GGG Gly	CCG Pro 70	GAG Glu	GAG Glu	CCG Pro	CTC Leu	CCC Pro 75	TAC Tyr	TCC Ser	CGG Arg	GCT Ala	TTC Phe 80	2	240
GGA Gly	GAG Glu	GGT Gly	GCG Ala	TCC Ser 85	GCG Ala	CGG Arg	CCG Pro	CGC Arg	TGC Cys 90	TGC Cys	AGG Arg	AAC Asn	GGC Gly	GGT Gly 95	ACC Thr	2	88
TGC Cys	GTG Val	CTG Leu	GGC Gly 100	AGC Ser	TTC Phe	TGC Cys	GTG Val	TGC Cys 105	CCG Pro	GCC Ala	CAC His	TTC Phe	ACC Thr 110	GGC Gly	CGC Arg	3	336
TAC Tyr	TGC Cys	GAG Glu 115	CAT His	GAC Asp	CAG Gln	AGG Arg	CGC Arg 120	AGT Ser	GAA Glu	TGC Cys	GGC Gly	GCC Ala 125	Leu	GAG Glu	CAC His		384
GGA Gly	GCC Ala 130	TGG Trp	ACC Thr	CTC Leu	CGC Arg	GCC Ala 135	Суѕ	CAC His	CTC Leu	TGC Cys	AGG Arg 140	Cys	ATC Ile	TTC Phe	GGG Gly		432
GCC Ala 145	·Leu	CAC His	TGC Cys	CTC Leu	CCC Pro	Leu	CAG Gln	ACG Thr	CCT Pro	GAC Asp 155	Arg	TGT Cys	GAC Asp	CCG Pro	AAA Lys 160	•	480
GAC Asp	TTC Phe	CTG Leu	GCC Ala	TCC Ser 165	His	GCT Ala	CAC His	GGG Gly	CCG Pro 170	Ser	GCC Ala	GGG Gly	GGC Gly	GCC Ala 175	CCC Pro		528
AGC Ser	CTG Leu	CTA Leu	CTC Leu 180	Leu	CTG Leu	CCC	TGC Cys	GCA Ala 185	Leu	CTG Leu	CAC His	C CGC Arg	CTC Leu 190	Leu	CGC Arg		576
CCG	GAT Asp	GCG Ala 195	Pro	GCG Ala	G CAC	CCI Pro	CGG Arg	Ser	CTG Leu	GTC Val	CCT Pro	TCC Ser 205	· Val	CTC Leu	CAG Gln		624
CGG	GAG Glu 210	Arg	G CGC	CCC Pro	TGC Cys	GGA Gly 215	/ Arg	G CCC	G GGA	CTI Leu	GG(Gl _y 22(/ His	CGC Arg	CTI Lev	TAA 1 *		672

FIG. 2

AMINO ACID ALIGNMENT BETWEEN CRIPTO AND CRIPTIN



Decoration 'Decoration #1': Shade (with solid black) residues that match the consensus named 'Consensus #1' exactly.